

Laboratory of Metabolomics

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Research topics:

- Interactions between *Fusarium* pathogens and cereal plants, both at vegetative and generative stage (seedlings, ears, kernels), as well as the influence of *Fusarium* metabolites on host plants,
- Identification of *Fusarium* species and chemotypes by molecular markers,
- Identification of resistance genes to leaf rust in wheat using PCR and validation of *Lr* gene markers,
- Mechanisms of the ability of *Trichoderma* and *Clonostachys* fungi to reduce the growth and mycotoxin production by *Fusarium* pathogens, molecular characterization of antagonistic fungi,
- Mycotoxin biosynthesis pathways – gene evolution and polymorphism in various *Fusarium* species,
- Role of secondary metabolites in the interaction between host-plant and fungal pathogens,
- Role of secondary metabolites in plant defense against abiotic stress conditions (acclimatization to drought and cold),
- Development of new methods of analysis of plant and fungal metabolites with use of high performance liquid chromatography (HPLC and UPLC) hyphenated with mass spectrometry.
- Profiling of secondary metabolites in plant tissues.

Major achievements:

- Development of a method for qualitative and quantitative assessment of the composition of individual prolamin protein classes in wheat kernels, facilitating the estimation of flour properties using high-performance capillary electrophoresis and rheological methods.
- Development of multiple PCR reaction to identify Glu-1 genes in wheat (*Triticum aestivum* L.) for the purpose of a prompt evaluation of genotypes with desirable technological traits.
- Clarification of quantitatively inherited resistance of triticale and barley to ear blight and seedling blight caused by *Fusarium culmorum* and *F. avenaceum*, including mycotoxins accumulation (deoxynivalenol and nivalenol) in kernels.
- Determination of toxin-formation ability of *Fusarium* pathogens infecting ears of corn (primarily *F. subglutinans*, *F. poae*, *F. proliferatum* and *F. graminearum*) and susceptibility of Polish maize hybrid varieties to infestation with these pathogens.
- Verification of the STS-PCR type DNA markers to identify resistance genes for brown rust of wheat (also by means of an interlaboratory ring test).
- Development of methods of analysis of various groups of plant and fungal secondary metabolites with use of HPLC/MS technique, in particular – flavone and isoflavone glycosides and glucuronates.
- Application of HPLC/MS to analysis of plant fitochelatins and identification of their oxidized and reduced forms present in variable amounts in different plant organs.
- Detection of changes in plant metabolomes in response to fungal infection and their differences that depend on physiological state of leaves of treated plants and way of infection.